

(2)

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RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/873,546

DATE: 06/21/2001

TIME: 17:30:08

Input Set : A:\05080-1.txt

Output Set: N:\CRF3\06212001\I873546.raw

ENTERED

3 <110> APPLICANT: Clark, Geoff
4 Ellis, Chad
5 Vos, Michelle
7 <120> TITLE OF INVENTION: Rig: Novel Ras-Related Gene
9 <130> FILE REFERENCE: NIH-05080
C--> 11 <140> CURRENT APPLICATION NUMBER: US/09/873,546
C--> 11 <141> CURRENT FILING DATE: 2001-06-04
11 <160> NUMBER OF SEQ ID NOS: 15
13 <170> SOFTWARE: PatentIn version 3.0
15 <210> SEQ ID NO: 1
16 <211> LENGTH: 4
17 <212> TYPE: PRT
18 <213> ORGANISM: Artificial Sequence
20 <220> FEATURE:
21 <223> OTHER INFORMATION: Synthetic
23 <220> FEATURE:
24 <221> NAME/KEY: SITE
25 <222> LOCATION: (4)..(4)
26 <223> OTHER INFORMATION: Xaa at this position can be any amino acid.
29 <400> SEQUENCE: 1
31 Cys Ala Ala Xaa
32 1
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35 <211> LENGTH: 33
36 <212> TYPE: DNA
37 <213> ORGANISM: Artificial Sequence
39 <220> FEATURE:
40 <223> OTHER INFORMATION: Synthetic
42 <400> SEQUENCE: 2
43 gggggatcca tgccggaaca gagtaacgat tac 33
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47 <211> LENGTH: 31
48 <212> TYPE: DNA
49 <213> ORGANISM: Artificial Sequence
51 <220> FEATURE:
52 <223> OTHER INFORMATION: Synthetic
54 <400> SEQUENCE: 3
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59 <211> LENGTH: 597
60 <212> TYPE: DNA
61 <213> ORGANISM: Homo sapiens
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66 agctcgctgg tgctgcgctt cgtgaagggc acgttccgcg acacctacat cccaccatc 120
68 gaggacacct accggcaggt gatcagctgc gacaagagcg tgtgcacgct gcagatcaca 180
70 gacaccaccg gcagccacca gttcccggcc atgcagcgcc tgtccatctc caagggccac 240

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72 gccttcatcc tgggtgttctc cgtcaccagc aagcagtcgc tggaggagct ggggcccatac 300
74 tacaagctca tcgtgcagat caagggcagc gtggaggaca tccccgtgat gctcgtgggc 360
76 aacaagtgcg atgagacgca gcgggaggtg gacacgcgcg aggcgcaggc ggtggcccaa 420
78 gagtgggaagt gcgctttcat ggagacctcg gccaaatga actacaacgt caaggagctc 480
80 ttccaggagc tgcctgacgt ggagacgcgc cggaacatga gcctcaacat cgacggcaag 540
82 cgctccggga agcagaagag gacagaccgc gtcaagggca aatgcaccct catgtga 597

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86 <211> LENGTH: 198

87 <212> TYPE: PRT

88 <213> ORGANISM: Homo sapiens

90 <400> SEQUENCE: 5

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93 1 5 10 15

95 Gly Val Gly Lys Ser Ser Leu Val Leu Arg Phe Val Lys Gly Thr Phe

96 20 25 30

98 Arg Asp Thr Tyr Ile Pro Thr Ile Glu Asp Thr Tyr Arg Gln Val Ile

99 35 40 45

101 Ser Cys Asp Lys Ser Val Cys Thr Leu Gln Ile Thr Asp Thr Thr Gly

102 50 55 60

104 Ser His Gln Phe Pro Ala Met Gln Arg Leu Ser Ile Ser Lys Gly His

105 65 70 75 80

107 Ala Phe Ile Leu Val Phe Ser Val Thr Ser Lys Gln Ser Leu Glu Glu

108 85 90 95

110 Leu Gly Pro Ile Tyr Lys Leu Ile Val Gln Ile Lys Gly Ser Val Glu

111 100 105 110

113 Asp Ile Pro Val Met Leu Val Gly Asn Lys Cys Asp Glu Thr Gln Arg

114 115 120 125

116 Glu Val Asp Thr Arg Glu Ala Gln Ala Val Ala Gln Glu Trp Lys Cys

117 130 135 140

119 Ala Phe Met Glu Thr Ser Ala Lys Met Asn Tyr Asn Val Lys Glu Leu

120 145 150 155 160

122 Phe Gln Glu Leu Leu Thr Leu Glu Thr Arg Asn Met Ser Leu Asn

123 165 170 175

125 Ile Asp Gly Lys Arg Ser Gly Lys Gln Lys Arg Thr Asp Arg Val Lys

126 180 185 190

128 Gly Lys Cys Thr Leu Met

129 195

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132 <211> LENGTH: 21

133 <212> TYPE: DNA

134 <213> ORGANISM: Artificial Sequence

136 <220> FEATURE:

137 <223> OTHER INFORMATION: Synthetic

139 <400> SEQUENCE: 6

140 gcgtgggcaa gaactcgctg g

21

143 <210> SEQ ID NO: 7

144 <211> LENGTH: 21

145 <212> TYPE: DNA

146 <213> ORGANISM: Artificial Sequence

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148 <220> FEATURE:
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 151 <400> SEQUENCE: 7
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 160 <220> FEATURE:
 161 <223> OTHER INFORMATION: Synthetic
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 168 Val Lys Gly Lys
 169 20
 171 <210> SEQ ID NO: 9
 172 <211> LENGTH: 229
 173 <212> TYPE: PRT
 174 <213> ORGANISM: Homo sapiens
 176 <400> SEQUENCE: 9
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 179 1 5 10 15
 181 Leu Arg Leu Leu Pro Ala Leu Leu Ile Leu Arg Ala Phe Lys Pro His
 182 20 25 30
 184 Arg Lys Ile Arg Asp Tyr Arg Val Val Val Val Gly Thr Ala Gly Val
 185 35 40 45
 187 Gly Lys Ser Thr Leu Leu His Lys Trp Ala Ser Gly Asn Phe Arg His
 188 50 55 60
 190 Glu Tyr Leu Pro Thr Ile Glu Asn Thr Tyr Cys Gln Leu Leu Gly Cys
 191 65 70 75 80
 193 Ser His Gly Val Leu Ser Leu His Ile Thr Asp Ser Lys Ser Gly Asp
 194 85 90 95
 196 Gly Asn Arg Ala Leu Gln Arg His Val Ile Ala Arg Gly His Ala Phe
 197 100 105 110
 199 Val Leu Val Tyr Ser Val Thr Lys Lys Glu Thr Leu Glu Glu Lys
 200 115 120 125
 202 Ala Phe Tyr Glu Leu Ile Cys Lys Ile Lys Gly Asn Asn Leu His Lys
 203 130 135 140
 205 Phe Pro Ile Val Leu Val Gly Asn Lys Ser Asp Asp Thr His Arg Glu
 206 145 150 155 160
 208 Val Ala Leu Asn Asp Gly Ala Thr Cys Ala Met Glu Trp Asn Cys Ala
 209 165 170 175
 211 Phe Met Glu Ile Ser Ala Lys Thr Asp Val Asn Val Gln Glu Leu Phe
 212 180 185 190
 214 His Met Leu Leu Asn Tyr Lys Lys Lys Pro Thr Thr Gly Leu Gln Glu
 215 195 200 205
 217 Pro Glu Lys Lys Ser Gln Met Pro Asn Thr Thr Glu Lys Leu Leu Asp
 218 210 215 220
 220 Lys Cys Ile Ile Met

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221 225
223 <210> SEQ ID NO: 10
224 <211> LENGTH: 206
225 <212> TYPE: PRT
226 <213> ORGANISM: Homo sapiens
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231 1 5 10 15
233 Val Ile Met Val Gly Ser Gly Gly Val Gly Lys Ser Ala Leu Thr Leu
234 20 25 30
236 Gln Phe Met Tyr Asp Glu Phe Val Glu Asp Tyr Glu Pro Thr Lys Ala
237 35 40 45
239 Asp Ser Tyr Arg Lys Lys Val Val Leu Asp Gly Glu Glu Val Gln Ile
240 50 55 60
242 Asp Ile Leu Asp Thr Ala Gly Gln Glu Asp Tyr Ala Ala Ile Arg Asp
243 65 70 75 80
245 Asn Tyr Phe Arg Ser Gly Glu Gly Phe Leu Cys Val Phe Ser Ile Thr
246 85 90 95
248 Glu Met Glu Ser Phe Ala Ala Thr Ala Asp Phe Arg Glu Gln Ile Leu
249 100 105 110
251 Arg Val Lys Glu Asp Glu Asn Val Pro Phe Leu Leu Val Gly Asn Lys
252 115 120 125
254 Ser Asp Leu Glu Asp Lys Arg Gln Val Ser Val Glu Glu Ala Lys Asn
255 130 135 140
257 Arg Ala Glu Gln Trp Asn Val Asn Tyr Val Glu Thr Ser Ala Lys Thr
258 145 150 155 160
260 Arg Ala Asn Val Asp Lys Val Phe Phe Asp Leu Met Arg Glu Ile Arg
261 165 170 175
263 Ala Arg Lys Met Glu Asp Ser Lys Glu Lys Asn Gly Lys Lys Lys Arg
264 180 185 190
266 Lys Ser Leu Ala Lys Arg Ile Arg Glu Arg Cys Cys Ile Leu
267 195 200 205
269 <210> SEQ ID NO: 11
270 <211> LENGTH: 184
271 <212> TYPE: PRT
272 <213> ORGANISM: Homo sapiens
274 <400> SEQUENCE: 11
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277 1 5 10 15
279 Ser Ala Leu Thr Val Gln Phe Val Gln Gly Ile Phe Val Glu Lys Tyr
280 20 25 30
282 Asp Pro Thr Ile Glu Asp Ser Tyr Arg Lys Gln Val Glu Val Asp Cys
283 35 40 45
285 Gln Gln Cys Met Leu Glu Ile Leu Asp Thr Ala Gly Thr Glu Gln Phe
286 50 55 60
288 Thr Ala Met Arg Asp Leu Tyr Met Lys Asn Gly Gln Gly Phe Ala Leu
289 65 70 75 80
291 Val Tyr Ser Ile Thr Ala Gln Ser Thr Phe Asn Asp Leu Gln Asp Leu
292 85 90 95

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294 Arg Glu Gln Ile Leu Arg Val Lys Asp Thr Glu Asp Val Pro Met Ile
295           100           105           110
297 Leu Val Gly Asn Lys Cys Asp Leu Glu Asp Glu Arg Val Val Gly Lys
298           115           120           125
300 Glu Gln Gly Gln Asn Leu Ala Arg Gln Trp Cys Asn Cys Ala Phe Leu
301           130           135           140
303 Glu Ser Ser Ala Lys Ser Lys Ile Asn Val Asn Glu Ile Phe Tyr Asp
304 145           150           155           160
306 Leu Val Arg Gln Ile Asn Arg Lys Thr Pro Val Glu Lys Lys Lys Pro
307           165           170           175
309 Lys Lys Lys Ser Cys Leu Leu Leu
310           180
312 <210> SEQ ID NO: 12
313 <211> LENGTH: 183
314 <212> TYPE: PRT
315 <213> ORGANISM: Homo sapiens
317 <400> SEQUENCE: 12
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320 1           5           10           15
322 Ser Ala Leu Thr Val Gln Phe Val Thr Gly Thr Phe Ile Glu Lys Tyr
323           20           25           30
325 Asp Pro Thr Ile Glu Asp Phe Tyr Arg Lys Glu Ile Glu Val Asp Ser
326           35           40           45
328 Ser Pro Ser Val Leu Glu Ile Leu Asp Thr Ala Gly Thr Glu Gln Phe
329           50           55           60
331 Ala Ser Met Arg Asp Leu Tyr Ile Lys Asn Gly Gln Gly Phe Ile Leu
332 65           70           75           80
334 Val Tyr Ser Leu Val Asn Gln Gln Ser Phe Gln Asp Ile Lys Pro Met
335           85           90           95
337 Arg Asp Gln Ile Ile Arg Val Lys Arg Tyr Glu Lys Val Pro Val Ile
338           100          105          110
340 Leu Val Gly Asn Lys Val Asp Leu Glu Ser Glu Arg Glu Val Ser Ser
341           115          120          125
343 Ser Glu Gly Arg Ala Leu Ala Glu Glu Trp Gly Cys Pro Phe Met Glu
344           130          135          140
346 Thr Ser Ala Lys Ser Lys Thr Met Val Asp Glu Leu Phe Ala Glu Ile
347 145           150           155           160
349 Val Arg Gln Met Asn Tyr Ala Ala Gln Pro Asp Lys Asp Asp Pro Cys
350           165           170           175
352 Cys Ser Ala Cys Asn Ile Gln
353           180
355 <210> SEQ ID NO: 13
356 <211> LENGTH: 189
357 <212> TYPE: PRT
358 <213> ORGANISM: Homo sapiens
360 <400> SEQUENCE: 13
362 Met Thr Glu Tyr Lys Leu Val Val Val Gly Ala Gly Gly Val Gly Lys
363 1           5           10           15
365 Ser Ala Leu Thr Ile Gln Leu Ile Gln Asn His Phe Val Asp Glu Tyr

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VERIFICATION SUMMARY

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Input Set : A:\05080-1.txt

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L:11 M:270 C: Current Application Number differs, Replaced Current Application No

L:11 M:271 C: Current Filing Date differs, Replaced Current Filing Date

L:31 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1